Pct/09

Errors Corrected by the STIC oms Branch

Serial Number: 09/857,581 ENTERED

CRF Processing Date: 2/19/2001

Edited by:

Verified by: (STIC staff

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line. BE
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☑ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: 55
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

"Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING DATE: 07/19/2001
PATENT APPLICATION: **US/09/857,581** TIME: 08:11:10

Input Set : **A:\Pto.amc**
Output Set: **N:\CRF3\07192001\I857581.raw**

P.5

3 <110> APPLICANT: E. I. du Pont de Nemours and Company
5 <120> TITLE OF INVENTION: Nucleic Acid Sequences Encoding Isoflavone Synthase
7 <130> FILE REFERENCE: BB1339 PCT
C--> 9 **<140> CURRENT APPLICATION NUMBER: US/09/857,581**
C--> 10 **<141> CURRENT FILING DATE: 2001-06-05**
12 <150> PRIOR APPLICATION NUMBER: 60/117,769
13 <151> PRIOR FILING DATE: 1999-01-27
15 <150> PRIOR APPLICATION NUMBER: 60/144,783
16 <151> PRIOR FILING DATE: 1999-07-20
18 <150> PRIOR APPLICATION NUMBER: 60/156,094
19 <151> PRIOR FILING DATE: 1999-09-24
21 <160> NUMBER OF SEQ ID NOS: 66
23 <170> SOFTWARE: Microsoft Office 97
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 1756
27 <212> TYPE: DNA
28 <213> ORGANISM: Glycine max
30 <400> SEQUENCE: 1
31 gtaattaacc tcactcaaac tcgggatcac agaaaccaac aacagttctt gcactgaggt 60
32 ttcacgatgt tgctggaact tgcacttggt ttgtttgtgt tagctttgtt tctgcacttg 120
33 cgtcccacac caagtgcaaa atcaaaagca cttcgccacc tcccaaaccc tccaagccca 180
34 aagcctcgtc ttcccttcat tggccacctt cacctcttaa aagataaact tctccactat 240
35 gcactcatcg atctctccaa aaagcatggc cccttattct ctctctcctt cggctccatg 300
36 ccaaccgtcg ttgcctccac ccctgagttg ttcaagctct tcctccaaac ccacgaggca 360
37 acttccttca acacaaggtt ccaaacctct gccataagac gcctcactta cgacaactct 420
38 gtggccatgg ttccattcgg accttactgg aagttcgtga ggaagctcat catgaacgac 480
39 cttctcaacg ccaccaccgt caacaagctc aggcctttga ggacccaaca gatccgcaag 540
40 ttccttaggg ttatggccca aagcgcagag gcccagaagc cccttgacgt caccgaggag 600
41 cttctcaaat ggaccaacag caccatctcc atgatgatgc tcggcgaggc tgaggagatc 660
42 agagacatcg ctcgcgaggt tcttaagatc ttcggcgaat acagcctcac tgacttcatc 720
43 tggcctttga agtatctcaa ggttggaaag tatgagaaga ggattgatga catcttgaac 780
44 aagttcgacc ctgtcgttga aagggtcatc aagaagcgcc gtgagatcgt cagaaggaga 840
45 aagaacggag aagttgttga gggcgaggcc agcggcgtct tcctcgacac tttgcttgaa 900
46 ttcgctgagg acgagaccat ggagatcaaa attaccaagg agcaaatcaa gggccttgtt 960
47 gtcgactttt tctctgcagg gacagattcc acagcggtgg caacagagtg ggcattggca 1020
48 gagctcatca acaatcccag ggtgttgcaa aaggctcgtg aggaggtcta cagtgttgtg 1080
49 ggcaaagata gactcgttga cgaagttgac actcaaaacc ttccttacat tagggccatt 1140
50 gtgaaggaga cattccgaat gcacccacca ctcccagtgg tcaaaagaaa gtgcacagaa 1200
51 gagtgtgaga ttaatgggta tgtgatccca gagggagcat tggttctttt caatgtttgg 1260
52 caagtaggaa gggaccccaa atactgggac agaccatcag aattccgtcc cgagaggttc 1320
53 ttagaaactg gtgctgaagg ggaagcaggg cctcttgatc ttaggggcca gcatttccaa 1380
54 ctcctcccat ttgggtctgg gaggagaatg tgccctggtg tcaatttggc tacttcagga 1440
55 atggcaacac ttcttgcatc tcttatccaa tgctttgacc tgcaagtgct gggccctcaa 1500
56 ggacaaatat tgaaaggtga tgatgccaaa gttagcatgg aagagagagc tggcctcaca 1560
57 gttccaaggg cacatagtct cgtttgtgtt ccacttgcaa ggatcggcgt tgcatctaaa 1620
58 ctcctttctt aattaagata atcatcatat acaatagtag tgtcttgcca tcgcagttgc 1680
59 tttttatgta ttcataatca tcatttcaat aaggtgtgac tggtacttaa tcaagtaatt 1740

RAW SEQUENCE LISTING DATE: 07/19/2001
PATENT APPLICATION: **US/09/857,581** TIME: 08:11:10

Input Set : **A:\Pto.amc**
Output Set: **N:\CRF3\07192001\I857581.raw**

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62 <210> SEQ ID NO: 2
63 <211> LENGTH: 521
64 <212> TYPE: PRT
65 <213> ORGANISM: Glycine max
67 <400> SEQUENCE: 2
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69   1               5                  10                  15
71 His Leu Arg Pro Thr Pro Ser Ala Lys Ser Lys Ala Leu Arg His Leu
72              20                  25                  30
74 Pro Asn Pro Pro Ser Pro Lys Pro Arg Leu Pro Phe Ile Gly His Leu
75          35                  40                  45
77 His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Leu Ile Asp Leu Ser
78      50                  55                  60
80 Lys Lys His Gly Pro Leu Phe Ser Leu Ser Phe Gly Ser Met Pro Thr
81  65                  70                  75                  80
83 Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln Thr His
84                  85                  90                  95
86 Glu Ala Thr Ser Phe Asn Thr Arg Phe Gln Thr Ser Ala Ile Arg Arg
87             100                 105                 110
89 Leu Thr Tyr Asp Asn Ser Val Ala Met Val Pro Phe Gly Pro Tyr Trp
90         115                 120                 125
92 Lys Phe Val Arg Lys Leu Ile Met Asn Asp Leu Leu Asn Ala Thr Thr
93     130                 135                 140
95 Val Asn Lys Leu Arg Pro Leu Arg Thr Gln Gln Ile Arg Lys Phe Leu
96 145                 150                 155                 160
98 Arg Val Met Ala Gln Ser Ala Glu Ala Gln Lys Pro Leu Asp Val Thr
99                 165                 170                 175
101 Glu Glu Leu Leu Lys Trp Thr Asn Ser Thr Ile Ser Met Met Met Leu
102             180                 185                 190
104 Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu Lys Ile
105         195                 200                 205
107 Phe Gly Glu Tyr Ser Leu Thr Asp Phe Ile Trp Pro Leu Lys Tyr Leu
108     210                 215                 220
110 Lys Val Gly Lys Tyr Glu Lys Arg Ile Asp Asp Ile Leu Asn Lys Phe
111 225                 230                 235                 240
113 Asp Pro Val Val Glu Arg Val Ile Lys Lys Arg Arg Glu Ile Val Arg
114                 245                 250                 255
116 Arg Arg Lys Asn Gly Glu Val Val Glu Gly Glu Ala Ser Gly Val Phe
117             260                 265                 270
119 Leu Asp Thr Leu Leu Glu Phe Ala Glu Asp Glu Thr Met Glu Ile Lys
120         275                 280                 285
122 Ile Thr Lys Glu Gln Ile Lys Gly Leu Val Val Asp Phe Phe Ser Ala
123     290                 295                 300
125 Gly Thr Asp Ser Thr Ala Val Ala Thr Glu Trp Ala Leu Ala Glu Leu
126 305                 310                 315                 320
128 Ile Asn Asn Pro Arg Val Leu Gln Lys Ala Arg Glu Glu Val Tyr Ser
129                 325                 330                 335
131 Val Val Gly Lys Asp Arg Leu Val Asp Glu Val Asp Thr Gln Asn Leu
```

RAW SEQUENCE LISTING DATE: 07/19/2001
PATENT APPLICATION: **US/09/857,581** TIME: 08:11:10

Input Set : **A:\Pto.amc**
Output Set: **N:\CRF3\07192001\I857581.raw**

```
132              340                 345                 350
134 Pro Tyr Ile Arg Ala Ile Val Lys Glu Thr Phe Arg Met His Pro Pro
135          355                 360                 365
137 Leu Pro Val Val Lys Arg Lys Cys Thr Glu Glu Cys Glu Ile Asn Gly
138      370                 375                 380
140 Tyr Val Ile Pro Glu Gly Ala Leu Val Leu Phe Asn Val Trp Gln Val
141 385                 390                 395                 400
143 Gly Arg Asp Pro Lys Tyr Trp Asp Arg Pro Ser Glu Phe Arg Pro Glu
144                 405                 410                 415
146 Arg Phe Leu Glu Thr Gly Ala Glu Gly Glu Ala Gly Pro Leu Asp Leu
147              420                 425                 430
149 Arg Gly Gln His Phe Gln Leu Leu Pro Phe Gly Ser Gly Arg Arg Met
150          435                 440                 445
152 Cys Pro Gly Val Asn Leu Ala Thr Ser Gly Met Ala Thr Leu Leu Ala
153      450                 455                 460
155 Ser Leu Ile Gln Cys Phe Asp Leu Gln Val Leu Gly Pro Gln Gly Gln
156 465                 470                 475                 480
158 Ile Leu Lys Gly Asp Asp Ala Lys Val Ser Met Glu Glu Arg Ala Gly
159                 485                 490                 495
161 Leu Thr Val Pro Arg Ala His Ser Leu Val Cys Val Pro Leu Ala Arg
162              500                 505                 510
164 Ile Gly Val Ala Ser Lys Leu Leu Ser
165          515                 520
167 <210> SEQ ID NO: 3
168 <211> LENGTH: 27
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
175 <400> SEQUENCE: 3
176 cgggatccat gcaaccggaa accgtcg                                          27
178 <210> SEQ ID NO: 4
179 <211> LENGTH: 32
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
186 <400> SEQUENCE: 4
187 ccggaattct caccaaacat cacggaggta tc                                    32
189 <210> SEQ ID NO: 5
190 <211> LENGTH: 47
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
197 <400> SEQUENCE: 5
198 tcaaggagaa aaaaccccgg atccatgttg ctggaacttg cacttgg                    47
200 <210> SEQ ID NO: 6
201 <211> LENGTH: 35
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RAW SEQUENCE LISTING DATE: 07/19/2001
PATENT APPLICATION: **US/09/857,581** TIME: 08:11:10

Input Set : **A:\Pto.amc**
Output Set: **N:\CRF3\07192001\I857581.raw**

202 <212> TYPE: DNA
203 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide
208 <400> SEQUENCE: 6
209 ggccagtgaa ttgtaatacg actcactata gggcg 35
211 <210> SEQ ID NO: 7
212 <211> LENGTH: 24
213 <212> TYPE: DNA
214 <213> ORGANISM: Artificial Sequence
216 <220> FEATURE:
217 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
219 <400> SEQUENCE: 7
220 aaaattagcc tcacaaaagc aaag 24
222 <210> SEQ ID NO: 8
223 <211> LENGTH: 27
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Description of Artificial Sequence:PCR primer
230 <400> SEQUENCE: 8
231 atataaggat tgatagttta tagtagg 27
233 <210> SEQ ID NO: 9
234 <211> LENGTH: 1824
235 <212> TYPE: DNA
236 <213> ORGANISM: Glycine max
238 <400> SEQUENCE: 9
239 ggaaaattag cctcacaaaa gcaaagatca aacaaaccaa ggacgagaac acgatgttgc 60
240 ttgaacttgc acttggttta ttggttttgg ctctgtttct gcacttgcgt cccacaccca 120
241 ctgcaaaatc aaaagcactt cgccatctcc caaacccacc aagcccaaag cctcgtcttc 180
242 ccttcatagg acaccttcat ctcttaaaag acaaacttct ccactacgca ctcatcgacc 240
243 tctccaaaaa acatggtccc ttattctctc tctactttgg ctccatgcca accgttgttg 300
244 cctccacacc agaattgttc aagctcttcc tccaaacgca cgaggcaact tccttcaaca 360
245 caaggttcca aacctcagcc ataagacgcc tcacctatga tagctcagtg gccatggttc 420
246 ccttcggacc ttactggaag ttcgtgagga agctcatcat gaacgacctt cccaacgcca 480
247 ccactgtaaa caagttgagg cctttgagga cccaacagac ccgcaagttc cttagggtta 540
248 tggcccaagg cgcagaggca cagaagcccc ttgacttgac cgaggagctt ctgaaatgga 600
249 ccaacagcac catctccatg atgatgctcg gcgaggctga ggagatcaga gacatcgctc 660
250 gcgaggttct taagatcttt ggcgaataca gcctcactga cttcatctgg ccattgaagc 720
251 atctcaaggt tggaaagtat gagaagagga tcgacgacat cttgaacaag ttcgaccctg 780
252 tcgttgaaag ggtcatcaag aagcgccgtg agatcgtgag gaggagaaag aacggagagg 840
253 ttgttgaggg tgaggtcagc ggggttttcc ttgacacttt gcttgaattc gctgaggatg 900
254 agaccatgga gatcaaaatc accaaggacc acatcgaggg tcttgttgtc gacttttttct 960
255 cggcaggaac agactccaca gcggtggcaa cagagtgggc attggcagaa ctcatcaaca 1020
256 atcctaaggt gttggaaaag gctcgtgagg aggtctacag tgttgtggga aaggacagac 1080
257 ttgtggacga agttgacact caaaaccttc cttacattag agcaatcgtg aaggagacat 1140
258 tccgcatgca cccgccactc ccagtggtca aaagaaagtg cacagaagag tgtgagatta 1200
259 atggatatgt gatcccagag ggagcattga ttctcttcaa tgtatggcaa gtaggaagag 1260
260 accccaaata ctgggacaga ccatcggagt tccgtcctga gaggttccta gagacagggg 1320

RAW SEQUENCE LISTING DATE: 07/19/2001
PATENT APPLICATION: US/09/857,581 TIME: 08:11:10

Input Set : A:\Pto.amc
Output Set: N:\CRF3\07192001\I857581.raw

```
261 ctgaagggga agcagggcct cttgatctta ggggacaaca ttttcaactt ctcccatttg 1380
262 ggtctgggag gagaatgtgc cctggagtca atctggctac ttcgggaatg gcaacacttc 1440
263 ttgcatctct tattcagtgc ttcgacttgc aagtgctggg tccacaagga cagatattga 1500
264 agggtggtga cgccaaagtt agcatggaag agagagccgg cctcactgtt ccaagggcac 1560
265 atagtcttgt ctgtgttcca cttgcaagga tcggcgttgc atctaaactc ctttcttaat 1620
266 taagatcatc atcatatata atatttactt tttgtgtgtt gataatcatc atttcaataa 1680
267 ggtctcgttc atctactttt tatgaagtat ataagccctt ccatgcacat tgtatcatct 1740
268 cccatttgtc ttcgtttgct acctaaggca atctttttt ttttagaatc acatcatcct 1800
269 actataaact atcaatcctt atat                                       1824
271 <210> SEQ ID NO: 10
272 <211> LENGTH: 521
273 <212> TYPE: PRT
274 <213> ORGANISM: Glycine max
276 <400> SEQUENCE: 10
277 Met Leu Leu Glu Leu Ala Leu Gly Leu Leu Val Leu Ala Leu Phe Leu
278   1               5                  10                  15
280 His Leu Arg Pro Thr Pro Thr Ala Lys Ser Lys Ala Leu Arg His Leu
281              20                  25                  30
283 Pro Asn Pro Pro Ser Pro Lys Pro Arg Leu Pro Phe Ile Gly His Leu
284          35                  40                  45
286 His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Leu Ile Asp Leu Ser
287      50                  55                  60
289 Lys Lys His Gly Pro Leu Phe Ser Leu Tyr Phe Gly Ser Met Pro Thr
290  65                  70                  75                  80
292 Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln Thr His
293                  85                  90                  95
295 Glu Ala Thr Ser Phe Asn Thr Arg Phe Gln Thr Ser Ala Ile Arg Arg
296             100                 105                 110
298 Leu Thr Tyr Asp Ser Ser Val Ala Met Val Pro Phe Gly Pro Tyr Trp
299         115                 120                 125
301 Lys Phe Val Arg Lys Leu Ile Met Asn Asp Leu Pro Asn Ala Thr Thr
302     130                 135                 140
304 Val Asn Lys Leu Arg Pro Leu Arg Thr Gln Gln Thr Arg Lys Phe Leu
305 145                 150                 155                 160
307 Arg Val Met Ala Gln Gly Ala Glu Ala Gln Lys Pro Leu Asp Leu Thr
308             165                 170                 175
310 Glu Glu Leu Leu Lys Trp Thr Asn Ser Thr Ile Ser Met Met Met Leu
311             180                 185                 190
313 Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu Lys Ile
314         195                 200                 205
316 Phe Gly Glu Tyr Ser Leu Thr Asp Phe Ile Trp Pro Leu Lys His Leu
317     210                 215                 220
319 Lys Val Gly Lys Tyr Glu Lys Arg Ile Asp Asp Ile Leu Asn Lys Phe
320 225                 230                 235                 240
322 Asp Pro Val Val Glu Arg Val Ile Lys Lys Arg Arg Glu Ile Val Arg
323                 245                 250                 255
325 Arg Arg Lys Asn Gly Glu Val Val Glu Gly Glu Val Ser Gly Val Phe
326             260                 265                 270
328 Leu Asp Thr Leu Leu Glu Phe Ala Glu Asp Glu Thr Met Glu Ile Lys
```

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

FYI

VERIFICATION SUMMARY DATE: 07/19/2001
PATENT APPLICATION: **US/09/857,581** TIME: 08:11:11

Input Set : **A:\Pto.amc**
Output Set: **N:\CRF3\07192001\I857581.raw**

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:3373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3385 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3427 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3436 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3445 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66

PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/857,581

DATE: 06/27/2001
TIME: 15:07:36

Does Not Comply
Corrected Diskette Needed

Input Set : A:\BB1339 PCT.Seq Listing.txt
Output Set: N:\CRF3\06272001\I857581.raw

```
      3 <110> APPLICANT: E. I. du Pont de Nemours and Company
      5 <120> TITLE OF INVENTION: Nucleic Acid Sequences Encoding Isoflavone Synthase
      7 <130> FILE REFERENCE: BB1339 PCT
C-->  9 <140> CURRENT APPLICATION NUMBER: US/09/857,581
C--> 10 <141> CURRENT FILING DATE: 2001-06-05
     12 <150> PRIOR APPLICATION NUMBER: 60/117,769
     13 <151> PRIOR FILING DATE: 1999-01-27
     15 <150> PRIOR APPLICATION NUMBER: 60/144,783
     16 <151> PRIOR FILING DATE: 1999-07-20
     18 <150> PRIOR APPLICATION NUMBER: 60/156,094
     19 <151> PRIOR FILING DATE: 1999-09-24
     21 <160> NUMBER OF SEQ ID NOS: 66
     23 <170> SOFTWARE: Microsoft Office 97
```

ERRORED SEQUENCES

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     2554 <210> SEQ ID NO: 55
     2555 <211> LENGTH: 499
     2556 <212> TYPE: PRT
     2557 <213> ORGANISM: Lupinus albus
E--> 2559 <400> SEQUENCE: 49 55
     2560 Phe Leu His Leu Arg Pro Thr Pro Thr Ala Lys Ser Lys Ala Leu Arg
     2561   1               5                  10                  15
     2563 His Leu Pro Asn Pro Pro Ser Pro Lys Pro Arg Leu Pro Phe Ile Gly
     2564              20                  25                  30
     2566 His Leu His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Leu Ile Asp
     2567          35                  40                  45
     2569 Leu Ser Lys Lys His Gly Pro Leu Phe Ser Leu Tyr Phe Gly Ser Met
     2570      50                  55                  60
     2572 Pro Thr Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln
     2573  65                  70                  75                  80
     2575 Thr His Glu Ala Thr Ser Phe Asn Thr Arg Phe Gln Thr Ser Ala Ile
     2576                  85                  90                  95
     2578 Arg Arg Leu Thr Tyr Asp Ser Ser Val Ala Arg Val Pro Phe Gly Pro
     2579              100                 105                 110
     2581 Tyr Trp Lys Phe Val Arg Lys Leu Ile Met Asn Asp Leu Leu Asn Ala
     2582          115                 120                 125
     2584 Thr Thr Val Asn Lys Leu Arg Pro Leu Arg Thr Gln Gln Ile Arg Lys
     2585      130                 135                 140
     2587 Phe Leu Arg Val Met Ala Gln Gly Ala Glu Ala Gln Lys Pro Leu Asp
     2588 145                 150                 155                 160
     2590 Leu Thr Glu Glu Leu Leu Lys Trp Thr Asn Ser Thr Ile Ser Met Met
     2591                 165                 170                 175
     2593 Met Leu Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu
     2594              180                 185                 190
     2596 Lys Ile Phe Gly Glu Tyr Ser Leu Thr Asp Phe Ile Trp Pro Leu Lys
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RAW SEQUENCE LISTING DATE: 06/27/2001
PATENT APPLICATION: **US/09/857,581** TIME: 15:07:37

Input Set : **A:\BB1339 PCT Seq Listing.txt**
Output Set: **N:\CRF3\06272001\I857581.raw**

```
2597          195                 200                 205
2599 His Leu Lys Val Gly Lys Tyr Glu Lys Arg Ile Asp Asp Ile Leu Asn
2600     210                 215                 220
2602 Lys Phe Asp Pro Val Val Glu Arg Val Ile Lys Lys Arg Arg Glu Ile
2603 225                 230                 235                 240
2605 Val Arg Arg Arg Lys Asn Gly Glu Val Val Glu Gly Glu Val Ser Gly
2606                 245                 250                 255
2608 Val Leu Leu Asp Thr Leu Leu Glu Phe Ala Glu Asp Glu Thr Met Glu
2609             260                 265                 270
2611 Ile Lys Ile Thr Lys Asp His Ile Lys Gly Leu Val Val Asp Phe Phe
2612         275                 280                 285
2614 Ser Ala Gly Thr Asp Ser Thr Ala Val Ala Thr Glu Trp Ala Leu Ala
2615     290                 295                 300
2617 Glu Leu Ile Asn Asn Pro Lys Val Leu Glu Arg Ala Arg Glu Glu Val
2618 305                 310                 315                 320
2620 Tyr Ser Val Val Gly Lys Asp Arg Leu Val Asp Glu Val Asp Thr Gln
2621                 325                 330                 335
2623 Asn Leu Pro Tyr Ile Arg Ala Ile Val Lys Glu Thr Phe Arg Met His
2624             340                 345                 350
2626 Pro Pro Leu Pro Val Val Lys Arg Lys Cys Thr Glu Glu Cys Glu Ile
2627         355                 360                 365
2629 Asn Gly Tyr Val Ile Pro Glu Gly Ala Leu Ile Leu Phe Asn Val Trp
2630     370                 375                 380
2632 Gln Val Gly Arg Asp Pro Lys Tyr Trp Asp Arg Pro Ser Glu Phe Arg
2633 385                 390                 395                 400
2635 Pro Glu Arg Phe Leu Glu Thr Glu Ala Glu Gly Glu Ala Arg Pro Leu
2636                 405                 410                 415
2638 Asp Leu Arg Gly Gln His Phe Gln Leu Leu Pro Phe Gly Ser Gly Arg
2639             420                 425                 430
2641 Arg Met Cys Pro Gly Val Ile Leu Ala Thr Ser Gly Met Ala Thr Leu
2642         435                 440                 445
2644 Leu Ala Ser Leu Ile Gln Cys Phe Asp Leu Gln Val Leu Gly Pro Gln
2645     450                 455                 460
2647 Gly Gln Ile Leu Lys Gly Gly Asp Ala Lys Val Ser Met Glu Glu Arg
2648 465                 470                 475                 480
2650 Ala Gly Leu Thr Val Pro Arg Ala His Ser Leu Val Cys Val Pro Leu
2651                 485                 490                 495
2653 Ala Arg Ile
```

VERIFICATION SUMMARY DATE: 06/27/2001
PATENT APPLICATION: **US/09/857,581** TIME: 15:07:38

Input Set : **A:\BB1339 PCT Seq Listing.txt**
Output Set: **N:\CRF3\06272001\I857581.raw**

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2559 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:55 differs:49
L:3373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3385 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3427 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3436 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3445 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:3463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66